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# Ectomycorrhiza Formation of *Tricholoma matsutake* on Mature Pine Root in Situ

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**Abstract:** We studied on the ectomycorrhiza formation of *Tricholoma matsutake* on mature pine (*Pinus densiflora*) root using matsutake solid spawn in situ. To find out suitable medium for the solid spawn, the matsutake mycelium growths on different media were compared. Among them, GYP medium (glucose 20g, yeast extract 5g, peptone 5g, KH<sub>2</sub>PO<sub>4</sub> 1g, CaCl<sub>2</sub> 50mg and vitamin B<sub>1</sub> 50mg) and GWR medium (glucose 20g, wheat bran 50g, rice bran 50g, KH<sub>2</sub>PO<sub>4</sub> 1g, CaCl<sub>2</sub> 50mg and vitamin B<sub>1</sub> 50mg) showed good proliferation of matsutake mycelium. The GYP and GWR media were used to make charcoal and vermiculite-soil spawns. The ergosterol amounts of charcoal and vermiculite-soil spawns were 1.53 and 1.61 times higher than that of the natural shiro, respectively. In spring, the roots of about 30-year-old red pine were shortened with unsterilized scissors. About twelve months later, the regenerated root tips from shortened points were inoculated with charcoal and vermiculite-soil spawns, respectively. After four months by the inoculation with charcoal and vermiculite-soil spawns, 55% and 50% of experimental plots inoculated with charcoal and vermiculite-soil spawns revealed dichotomous root tips characteristic to ectomycorrhiza, respectively. The nested PCR amplification results confirmed that the dichotomous root tips were colonised by matsutake spawn. Our findings demonstrated that the ectomycorrhiza of *T. matsutake* could be formed on the mature pine root in situ by using matsutake solid spawns.

**Keywords:** *Tricholoma matsutake*, Ectomycorrhiza, Solid Spawn, Inoculation, *Pinus densiflora*, Nested PCR

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## 1. Introduction

*Tricholoma matsutake*, a member of the family Tricholomataceae, order Agaricales, has been reported from Korean peninsula, China, Japan and Bhutan, meanwhile *Tricholoma matsutake* in Korean peninsula, Japan, and northeastern China are closely related, but distinct from those in southwestern China and Bhutan [1]. In DPR Korea, the fruiting bodies of *T. matsutake* occur in almost areas including Mt. Myohyang, Mt. Kumgang, Buryong, Yangdok, Myongchen, Kimchaek, Kilju, Rason, Chosan etc.

*T. matsutake* usually occurs in Japanese red pine (*Pinus densiflora*) forests and also under trees of *Pinus thunbergii*, *Pinus pumila*, *Tsuga diversifolia*, *Tsuga sieboldii*, and *Abies*

*veitchii* [2].

Matsutakes appear on nutrient-poor soils, such as granite, chert, and sandstone, which are acidic (pH 4.5-5.0) and develops special structure called “shiros” in symbiotic relation with host root [3]. Shiros are mycelial aggregations that develop in association with the roots of host trees and soil particles in well drained and nutrient-poor forest soil, from which matsutake generates. The fruiting bodies of *T. matsutake* occur on the zone of around 25 cm inwards from the growing tip of the shiro, which is called an “active mycorrhizal zone” [4]. Shiros expand at a rate of 10~15 cm per year or 17 cm per year [5].

The matsutake has been known as one of the most expensive edible and medicinal mushroom in many countries



#### 2.4. Spawn Culture of Matsutake Isolates

Charcoal substrates with ca. 5mm diameter were treated with hydrochloric acid to make its pH to 5.5. Then the GYP medium was added into charcoal substrate up to 35% humidity, and the matsutake isolate was inoculated into charcoal substrate and incubated at 23°C for 40 ~ 50 days to make charcoal spawn. A vermiculite-soil spawn was made as follows: GWR medium was added into vermiculite-soil substrate (5: 5, v/v) up to 35% humidity, and the substrate was inoculated by the charcoal spawn and incubated at 23°C for 60 days. Ergosterol amounts of matsutake spawns and natural shiro were measured by the method of Zhao et al. [21].

#### 2.5. Inoculation of *T. matsutake* Spawn on Mature Pine Root

In spring, the roots (ca. 3~5mm in diameter) of about 30-year-old pines were shortened with scissors in 10~20cm depth of soil, and buried with fallen leaves and soil. About twelve months later, the regenerated roots from shortened points were sterilized with 75% alcohol and inoculated with charcoal and vermiculite-soil spawns, respectively. For the prevention of any native EM fungi, the inoculation portions were wrapped up with the riversand or 20cm-depth mineral soil and covered with a perforated vinyl sheet.

Inoculation of matsutake spawns were carried out in spring when soil temperature was about 10°C and the soil native microbes did not flourish.

#### 2.6. Morphological and Molecular Analysis of Ectomycorrhiza Formation by Matsutake Spawns

Inoculated root portions were excavated cautiously four months later, and primarily observed if there were dichotomic root tips and matsutake odor from root. Dichotomic root tips with matsutake odor were collected and observed under a dissecting microscope.

A nested PCR amplification was carried out on DNA extracts (CTAB method) from samples (about 15mg). Amplification using fungal-specific primers ITS1/ITS4 [22] was followed by reamplification using matsutake-specific primers TmF/TmR [20]. The first amplification with primers ITS1 and ITS4 was carried out in a 25 µl reaction solution including 2.5 µl 10× PCR buffer, 200 µM of each dNTP, 1 µM of each primer, 1.5U Taq polymerase and 1 µl template DNA. Reactions were run in a C1000 Touch™ Thermal Cycler. The thermal cycling was followed by an initial denaturation at 94°C for 10 min, 35 cycles of denaturation at 94°C for 40s, annealing at 55°C for 40s and extension at 72°C for 1min, and then a final extension at 72°C for 10 min. The product of the first amplification was diluted 10 times with sterilized deionized water and 1.0 µl of the diluted solution was used as the template for the nested PCR. Conditions for the nested PCR using primers TmF and TmR were similar to the first PCR, except for an initial denaturation at 95°C for 10 min and denaturation at 95°C for

40s. The PCR products were detected by electrophoresis in 0.8% (w/v) agarose gel supplemented with ethidium bromide. The DNA was visualized and photographed over a UV light source. A DM 2 000 molecular weight marker (CWBIO, China) was used as the size standard.

#### 2.7. Statistical Analysis

Statistical analysis was carried out in SPSS 22.0. Significant differences (mean ± SE) among treatments were compared by Tukey's HSD test at  $P < 0.05$ .

### 3. Results

#### 3.1. *T. matsutake* Isolates

Morphological characteristics of isolates on slants of Hamada's agar medium showed the typical characters of *T. matsutake* (Figure 1) and molecular analysis using matsutake-specific primers TmF/TmR also proved the isolates were *T. matsutake* (Figure 6).

#### 3.2. Selection of Optimum Medium

For the selection of optimum medium, matsutake isolates were grown in 11 different media including Hamada's agar media. Among them, GYP and GWR media showed good proliferation of matsutake mycelium with day growth rate of  $0.46 \pm 0.03$ ,  $0.53 \pm 0.01$  mm/d, respectively (Figure 2). The growth rates of *T. matsutake* on GYP and GWR media were 1.27 and 1.47 times higher than that on Hamada medium known as common matsutake medium.

#### 3.3. Spawn Culture of Matsutake Isolates

The mycelial density of charcoal spawn and vermiculite-soil spawns were higher than that of the natural shiro and all the spawns also had the odor of *T. matsutake* (Table 3, Figure 3).

The ergosterol amounts of charcoal and vermiculite-soil spawns were  $110.5 \pm 4.3$ ,  $116.1 \pm 5.8$  µg/g and 1.53, 1.61 times higher than that ( $72.1 \pm 3.2$  µg/g) of the natural shiro, respectively (Figure 4).

In spring, the roots (ca. 3~5mm in diameter) of about 30-year-old red pines were shortened. About twelve months later, the regenerated root tips from shortened points (Figure 5. A, B) were inoculated by spawns. Inoculation was carried out in May when the soil temperature was about 9~13°C.

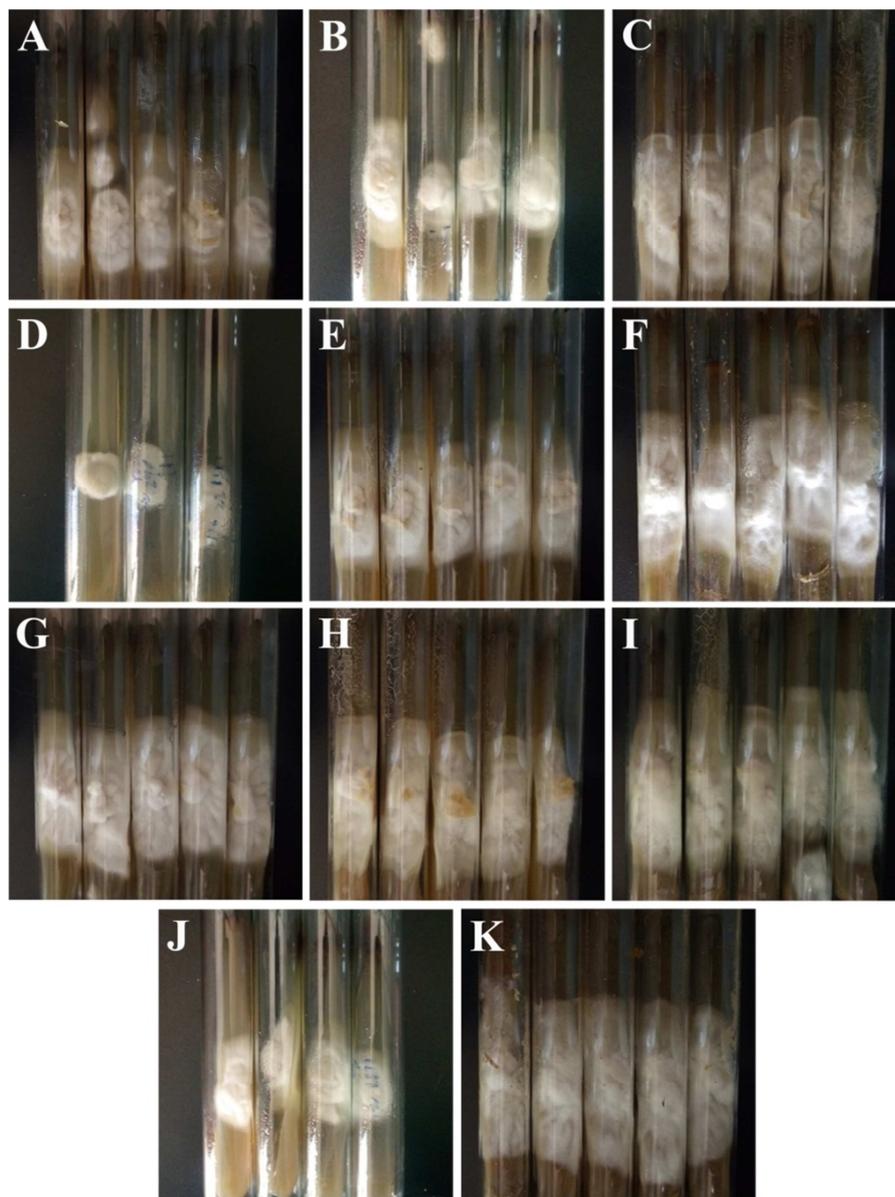
Four months later after inoculation, the inoculated portions were examined. The dichotomic root tips, characteristic to ectomycorrhiza, were formed in the inoculated root portions (Figure 5C, D, E).

DNA analysis confirmed the presence of *T. matsutake* in matsutake inoculated roots (Figure 6). No matsutake-specific bands were amplified from root samples of non-inoculated root portions as control. More than 50% of samples showed the dichotomic root tips characteristic to putative matsutake mycorrhiza, and the nested PCR results from DNA extracts

of any sample were also positive (Table 4).

We found that charcoal and vermiculite-soil spawns could be used as matsutake inoculum on the mature pine roots, but

we consider the results must be confirmed in the future with the development of matsutake fruiting body.



**Figure 1.** *T. matsutake* grown on different kinds of media. A: HA, B: PDA, C: GYP, D: YPP, E: YPR, F: YR, G: GY, H: YRB, I: YP, J: GPP, K: GWR.

**Table 3.** Culture properties of spawns.

Spawns	Germination (d)	Spawn running (d)	Hyphal density <sup>1)</sup>	Odor <sup>2)</sup>
Charcoal spawn	10	50	+	+
Vermiculite-soil spawn	7	45	+	+

<sup>1)</sup>: + higher than natural shiro. <sup>2)</sup>: + matsutake odor.

**Table 4.** Examination of inoculated sites in 2021 yr.

Sites (spawn)	Inoculation		Examination			
	Date	Number of trial	Date	Number of examination	Number of dichotomic root-site	Molecular analysis
HaeYeon-ri (vermiculite-soil spawn)	2021.5	30	2021.9	14	7	+
	2021.5	15	2021.9	8	4	+
GeSang-ri (charcoal spawn)	2021.5	40	2021.9	20	11	+

+ positive in nested PCR amplification

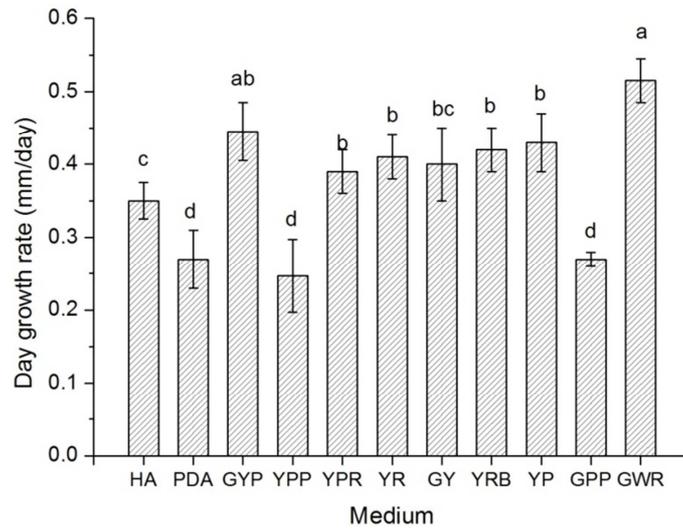


Figure 2. Effects of various media on mycelial growth of *Tricholoma matsutake*. Bars without shared letters indicate significant difference at  $P < 0.05$ .

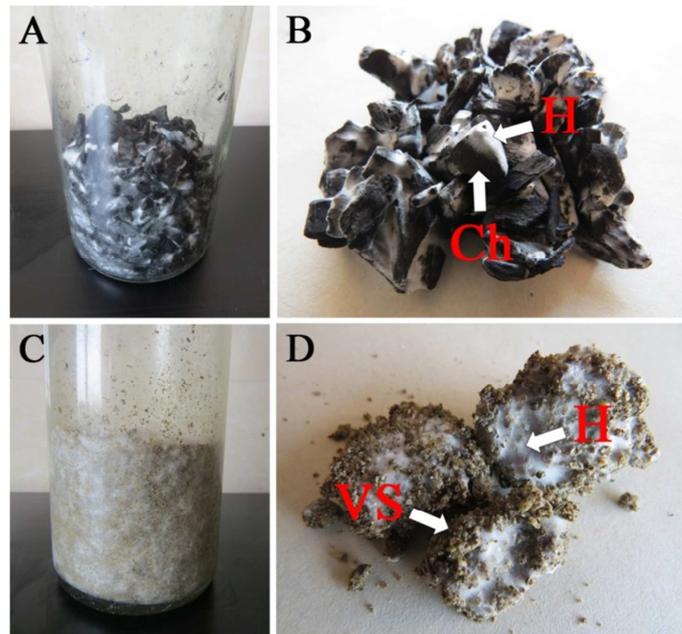


Figure 3. Charcoal spawn and vermiculite-soil spawn. A: Charcoal spawn in bottle, B: Charcoal spawn used in inoculation, C: Vermiculite-soil spawn in bottle, D: Vermiculite-soil spawn used in inoculation. Abbreviations, H: matsutake hyphae, Ch: charcoal, VS: vermiculite-soil.

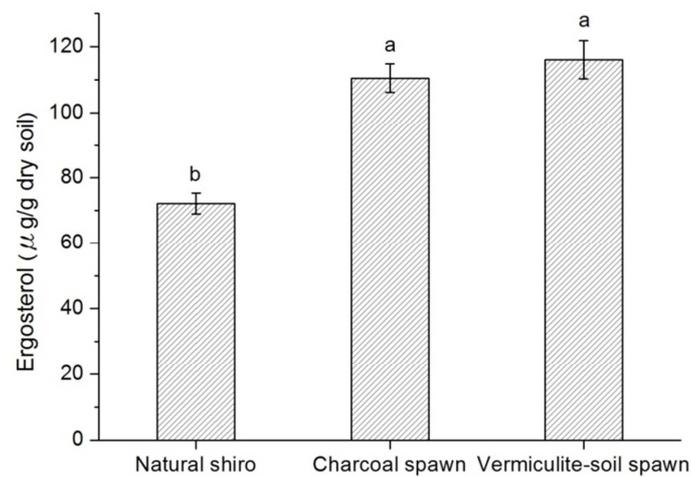
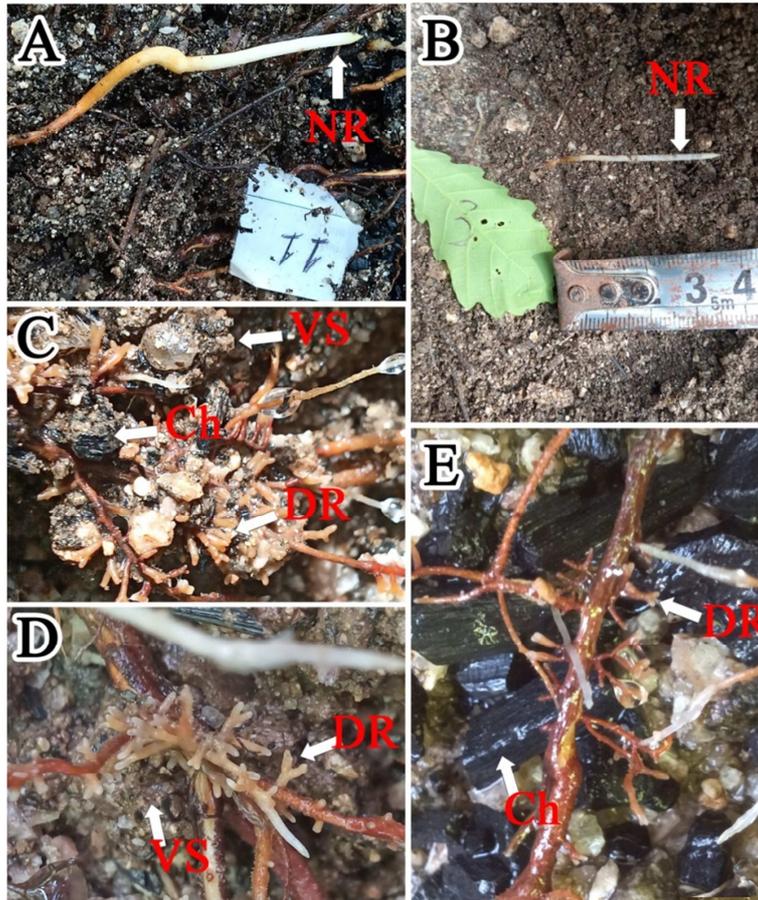
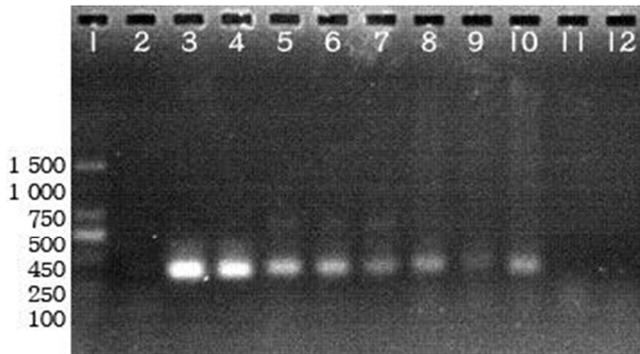


Figure 4. The amounts of ergosterol in natural shiro and spawns. Bars without shared letters indicate significant difference at  $P < 0.05$ .



**Figure 5.** New generated root tips (NR) and dichotomic root tips (DR) observed in inoculated sites. *A* and *B*: New generated root tips from shortened points, *C*: Dichotomic root tips in charcoal (Ch) and vermiculite-soil spawn (VS) inoculated sites, *D*: Dichotomic root tips in vermiculite-soil spawn inoculated sites, *E*: Dichotomic root tips in charcoal spawn inoculated sites.



**Figure 6.** Nested PCR on DNA extracts from inoculated root samples. Second step reamplification of the samples, using the matsutake specific primer pair TmF/TmR. (1: DNA maker, 2: Negative control, 3: Fruiting body of *T. matsutake*, 4: *T. matsutake* isolates, 5: Natural shiwo mycelium, 6: Natural shiwo root, 7 and 8: Charcoal spawn inoculated root, 9, 10: Vermiculite-soil inoculated root, 11 and 12: control).

### 4. Conclusion

*T. matsutake* is a symbiotic ectomycorrhizal fungus which is mainly produced on root of pine tree [6]. The growth of *T. matsutake*, which uses symbiotic nutrition, is very slow on artificial media [18], so that it is necessary to examine the effects of various nutrients on the matsutake mycelial growth

in order to produce matsutake spawns. Although many nutrients have good effects on matsutake mycelial growth [23], we found that the media including yeast extracts, wheat bran, rice bran, and vitamin B<sub>1</sub> gave the best mycelial growth for preparation of matsutake spawns.

Many studies were carried out to form artificial matsutake ectomycorrhiza on pine roots [17, 18]. Yamada *et al.* inoculated matsutake isolates on 1-wk-old seedlings, and demonstrated that the fungus formed a sheath and Hartig net on the pine lateral roots to form successfully matsutake ectomycorrhiza during three months after inoculation [10]. This result indicates that cultured *T. matsutake* mycelium can form true ectomycorrhizas with *P. densiflora* seedlings. Especially, Guerin-Laguette *et al.* regenerated mycorrhiza-free short roots in 50-year-old *Pinus densiflora* trees and inoculated mycelial pieces of *T. matsutake* on mycorrhiza-free short roots [17]. After four-and-a-half months, the matsutake-inoculated-roots were colonized by matsutake mycelia. Although this study was limited to a single tree, the results clearly confirm the symbiotic infection of mature *P. densiflora* roots by matsutake isolates. In order to produce fruiting bodies of *T. matsutake*, mass liquid inoculum of matsutake isolates were inoculated on pine forest ground and after one year of inoculation, no external difference was observed between the artificially inoculated mycelia and the

naturally existing mycelia of *T. matsutake* [18].

In their study, the results of artificial *T. matsutake* development have not been reported but they suggested that using liquid inoculum of *T. matsutake*, used in their research, may be new technique to obtain artificial *T. matsutake*. *T. matsutake* is an obligate fungus that forms a symbiotic association with the roots of living pine trees and receives nutrients from them [7-9], so that it is very important to produce spawn to form artificial ectomycorrhiza on the roots of mature pine for the cultivation of *T. matsutake* in pine forest. Previous studies on ectomycorrhiza formation of matsutake in pine root mostly used matsutake mycelia on agar media or liquid medium as inoculum [10, 17, 18].

We predicted that the inoculum with nutrient-poor substrates such as charcoal, vermiculite and mineral soil would be favorable to mycorrhiza formation in situ, because spawn with nutrient-rich substrates could be decayed in the soil for months to damage the host roots inoculated with matsutake spawn. Our results indicate that *T. matsutake* spawns cultivated in charcoal or vermiculite-soil can form ectomycorrhizas with mature pine roots in situ.

## References

- [1] BAO D., KOIKE A., YAO F., YAMANAKA K., AIMI T., AND KITAMOTO Y. 2007. Analyses of the genetic diversity of matsutake isolates collected from different ecological environments in Asia. *Journal of Wood Science*. 53: 344-350.
- [2] ENDO N., DOKMAI P., SUWANNASAI N., PHOSRI C., HORIMAI Y., HIRAI N., et al. 2015. Ectomycorrhization of *Tricholoma matsutake* with *Abies veitchii* AND *Tsuga diversifolia* in the subalpine forests of Japan. *Mycoscience*. 56: 402-412.
- [3] VAARIO L. M., SAVONEN E. M., PELTONIEMI M., MIYAZAWA T., PULKKINEN P., AND SARJALAT. 2015. Fruiting pattern of *Tricholoma matsutake* in southern Finland. *Scandinavian Journal of Forest Research*. 30: 259-265.
- [4] OHARA H., AND HAMADA M. 1967. Disappearance of bacteria from the zone of active mycorrhizas in *Tricholoma matsutake* (S. Ito et Imai) singer. *Nature*. 213: 528-529.
- [5] NARIMATSU M., KOIWA T., MASAKI T., SAKAMOTO Y., OHMORI H., AND TAWARAYA K. 2015. Relationship between climate, expansion rate, and fruiting in fairy rings ('shiro') of an ectomycorrhizal fungus *Tricholoma matsutake* in a *Pinus densiflora* forest. *Fungal Ecology*. 15: 18-28.
- [6] TAKASHI Y., AKIYOSHI Y., AND HITOSHI F. 2020. Advances in the cultivation of the highly-prized ectomycorrhizal mushroom *Tricholoma matsutake*. *Mycoscience*. 61: 49-57.
- [7] KUSUDA M., UEDA M., MIYATAKE K., AND TERASHITA T. 2008. Characterization of the carbohydrase productions of an ectomycorrhizal fungus, *Tricholoma matsutake*. *Mycoscience*. 49: 291-297. <https://doi.org/10.1007/S10267-008-0423-7>.
- [8] KUSUDA M., UEDA M., NAKAZAWA M., MIYATAKE K., AND YAMANAKA K. 2006. Detection of b-glucosidase as saprotrophic ability from an ectomycorrhizal mushroom, *Tricholoma matsutake*. *Mycoscience*. 47: 184e189.
- [9] SHIMOKAWA T., YAMAGUCHI M., AND MURATA H. 2017. Agar plate assays using dye linked substrates differentiate members of *Tricholoma* sect. *caligata*, ectomycorrhizal symbionts represented by *Tricholoma matsutake*. *Mycoscience*. 58: 432-437.
- [10] YAMADA A., MAEDA K., AND OHMASA M. 1999a. Ectomycorrhiza formation of *Tricholoma matsutake* isolates on seedlings of *Pinus densiflora* in vitro. *Mycoscience*. 40: 455-463.
- [11] GUERIN-LAGUETTE A., VAARIO L. M., GILLW M., LAPEYRIE F., MATSUSHITA N., AND SUZUKI K. 2000. Rapid in vitro ectomycorrhizal infection on *Pinus densiflora* roots by *Tricholoma matsutake*. *Mycoscience*. 41: 389-393.
- [12] VAARIO L. M., GUERIN-LAGUETTE A., GILL W. M., LAPEYRIE F., AND SUZUKI K. 2000. Only two weeks are required for *Tricholoma matsutake* to differentiate ectomycorrhizal Hartig net structures in roots of *Pinus densiflora* seedlings cultivated on artificial substrate. *Journal of Forest Research*. 5: 293-297.
- [13] YAMADA A., KANEKAWA S., AND OHMASA M. 1999b. Ectomycorrhiza formation of *Tricholoma matsutake* on *Pinus densiflora*. *Mycoscience*. 40: 193-198. <https://doi.org/10.1007/BF02464298>.
- [14] GUERIN-LAGUETTE A., VAARIO L. M., MATSUSHITA N., SHINDO K., SUZUKI K., AND LAPEYRIE F. 2003. Growth stimulation of a Shiro-like, mycorrhiza forming, mycelium of *Tricholoma matsutake* on solid substrates by non-ionic surfactants or vegetable oil. *Mycological Progress*. 2: 37-44.
- [15] KA K. H., KIM H. S., HUR T. C., PARK H., JEON S. M., RYOO R., et al. 2018. Analysis of environment and production of *Tricholoma matsutake* in matsutake-infected pine trees (in Korean). *Korean Journal of Mycology*. 46: 34-42.
- [16] KAREKI K., AND KAWAKAMI Y. 1985. Artificial formation of Shiro (fungus colony) by planting the pine saplings infected with *Tricholoma matsutake* (Ito et Imai) Sing. *Bulletin of the Hiroshima Prefectural Forest Experiment Station*. 20: 13-23 (in Japanese).
- [17] GUERIN-LAGUETTE A., MATSUSHITA N., LAPEYRIE F., SHINDO K., AND SUZUKI K. 2005. Successful inoculation of mature pine with *Tricholoma matsutake*. *Mycorrhiza*. 15: 301-305. <https://doi.org/10.1007/s00572-009-0286-6>.
- [18] LEE W. H., HAN S. K., KIM B. S., BHUSHAN S, LEES Y., KOC S., SUNG G. H., AND SUNG J. M. 2007. Proliferation of *Tricholoma matsutake* Mycelial Mats in Pine Forest Using Mass Liquid Inoculum. *Mycobiology*. 35 (2): 54-61.
- [19] HAMADA M. 1964. Method of pure culture of the mycelium of *Armillaria matsutake* and of its allies. In: *The Matsutake Research Association (Ed.), Matsutake (Tricholoma matsutake Singer)-Its fundamental studies and economic production of the fruit-body (pp. 97-100)*. Kyoto: Nakanishi printing (in Japanese).
- [20] KIKUCHI K., MATSUSHITA N., GUERIN-LAGUETTE A., OHTA A. AND SUZUKI K. 2000. Detection of *Tricholoma matsutake* by specific primers. *Mycol. Res*. 104: 1427-1430.
- [21] ZHAO X. R., LINA Q., AND BROOKES P. C. 2005. Does soil ergosterol concentration provide a reliable estimate of soil fungal biomass? *Soil biology and biochemistry*. 37: 311-315.

- [22] WHITE T. J., BRUNS T., LEE S., AND TAYLOR J. 1990. Amplification and direct sequencing of fungal ribosomal RNA genes for phylogenetics. P 315–322 in Innis MA, Gelfand DH, Snisky JJ, White TJ (eds) PCR protocols. A guide to methods and applications. Academic, San Diego, CA.
- [23] Kawai M., IMAJI A., YAMADA A., AND KINOSHITA A. 2018. Shiro formation and fruit body flashing by inoculation of *Tricholoma bakamatsutake* mycelia in the forest. P. 25 in Abstracts of the annual meeting of the mycological Society of Japan. [https://doi.org/10.11556/msj7abst.62.0\\_25](https://doi.org/10.11556/msj7abst.62.0_25).